AMENDMENTS TO THE CLAIMS

1-112. (Cancelled)

- 113. (Currently Amended) An inhibitor for A method for inhibiting an interaction between a protein that interacts with a c-Jun protein and the c-Jun protein, which comprises adding, to a system in which an interaction between the protein and the c-Jun protein occurs, a protein of the following (a) or (b): as an active ingredient:
- (a) a protein comprising any one of the amino acid sequences of SEQ ID NOS: 1 to 69, SEQ ID NOS: 70 to 87, SEQ ID NOS: 88 to 94, SEQ ID NOS: 95 to 99, SEQ ID NOS: 100 to 104, SEQ ID NOS: 105 to 108, SEQ ID NOS: 109 to 111, SEQ ID NOS: 112 and 113, SEQ ID NOS: 114 and 115, SEQ ID NOS: 116 and 117, SEQ ID NOS: 118 and 119, SEQ ID NOS: 120 and 121, SEQ ID NOS: 122 and 123, and SEQ ID NOS: 124 and 125,
- (b) a protein that comprises any one of the amino acid sequences of SEQ ID NOS: 1 to 69, SEQ ID NOS: 70 to 87, SEQ ID NOS: 88 to 94, SEQ ID NOS: 95 to 99, SEQ ID NOS: 100 to 104, SEQ ID NOS: 105 to 108, SEQ ID NOS: 109 to 111, SEQ ID NOS: 112 and 113, SEQ ID NOS: 114 and 115, SEQ ID NOS: 116 and 117, SEQ ID NOS: 118 and 119, SEQ ID NOS: 120 and 121, SEQ ID NOS: 122 and 123, and SEQ ID NOS: 124 and 125, including deletion, substitution or addition of one or several amino acid residues and interacts with the c-Jun protein.
- 114. (Currently Amended) The methodinhibitor according to claim 113, wherein the protein as the active ingredient of the following (a) or (b) comprises any one of the amino acid sequences of SEQ ID NOS: 1 to 69, SEQ ID NOS: 70 to 87, SEQ ID NOS: 88 to 94, SEQ ID NOS: 95 to 99, SEQ ID NOS: 100 to 104, SEQ ID NOS: 105 to 108, SEQ ID NOS: 109 to 111, SEQ ID NOS: 112 and 113, SEQ ID NOS: 114 and 115, SEQ ID NOS: 116 and 117, SEQ ID NOS: 118 and 119, SEQ ID NOS: 120 and 121, SEQ ID NOS: 122 and 123, and SEQ ID NOS: 124 and 125.
- 115. (Currently Amended) The methodinhibitor according to claim 114, wherein the protein of the following (a) or (b) is a protein translated from a nucleic acid of the following (a) or (b):

- (a) a nucleic acid comprising any one of the nucleotide sequences of SEQ ID NOS: 126 to 199, SEQ ID NOS: 200 to 217, SEQ ID NOS: 218 to 224, SEQ ID NOS: 225 to 229, SEQ ID NOS: 230 to 234, SEQ ID NOS: 235 to 238, SEQ ID NOS: 239 to 241, SEQ ID NOS: 242 and 243, SEQ ID NOS: 244 and 245, SEQ ID NOS: 246 and 247, SEQ ID NOS: 248 and 249, SEQ ID NOS: 250 and 251, SEQ ID NOS: 252 and 253, and SEQ ID NOS: 254 and 255, (b) a nucleic acid that hybridizes with a nucleic acid comprising any one of the nucleotide sequences of SEQ ID NOS: 126 to 199, SEQ ID NOS: 200 to 217, SEQ ID NOS: 218 to 224, SEQ ID NOS: 225 to 229, SEQ ID NOS: 230 to 234, SEQ ID NOS: 235 to 238, SEQ ID NOS: 239 to 241, SEQ ID NOS: 242 and 243, SEQ ID NOS: 244 and 245, SEQ ID NOS: 246 and 247, SEQ ID NOS: 248 and 249, SEQ ID NOS: 250 and 251, SEQ ID NOS: 252 and 253, and SEQ ID NOS: 254 and 255, under a stringent condition and encodes a protein that interacts with the c-Jun protein.
- 116. (Currently Amended) The methodinhibitor according to claim 115, wherein the nucleic acid comprises any one of the nucleotide sequences of SEQ ID NOS: 126 to 199, SEQ ID NOS: 200 to 217, SEQ ID NOS: 218 to 224, SEQ ID NOS: 225 to 229, SEQ ID NOS: 230 to 234, SEQ ID NOS: 235 to 238, SEQ ID NOS: 239 to 241, SEQ ID NOS: 242 and 243, SEQ ID NOS: 244 and 245, SEQ ID NOS: 246 and 247, SEQ ID NOS: 248 and 249, SEQ ID NOS: 250 and 251, SEQ ID NOS: 252 and 253, and SEQ ID NOS: 254 and 255.
- 117. (Previously Presented) A method for detecting an interaction between a bait and a prey, which comprises bringing the bait and the prey into contact and detecting a complex formed by the contact, wherein the bait is a protein of the following (a) or (b) or a protein translated from a nucleic acid of the following (a') or (b'):
- (a) a protein comprising any one of the amino acid sequences of SEQ ID NOS: 1 to 69, SEQ ID NOS: 70 to 87, SEQ ID NOS: 88 to 94, SEQ ID NOS: 95 to 99, SEQ ID NOS: 100 to 104, SEQ ID NOS: 105 to 108, SEQ ID NOS: 109 to 111, SEQ ID NOS: 112 and 113, SEQ ID NOS: 114 and 115, SEQ ID NOS: 116 and 117, SEQ ID NOS: 118 and 119, SEQ ID NOS: 120 and 121, SEQ ID NOS: 122 and 123, and SEQ ID NOS: 124 and 125,
- (b) a protein that comprises any one of the amino acid sequences of SEQ ID NOS: 1 to 69, SEQ ID NOS: 70 to 87, SEQ ID NOS: 88 to 94, SEQ ID NOS: 95 to 99, SEQ ID NOS: 100 to 104,

SEQ ID NOS: 105 to 108, SEQ ID NOS: 109 to 111, SEQ ID NOS: 112 and 113, SEQ ID NOS: 114 and 115, SEQ ID NOS: 116 and 117, SEQ ID NOS: 118 and 119, SEQ ID NOS: 120 and 121, SEQ ID NOS: 122 and 123, and SEQ ID NOS: 124 and 125, including deletion, substitution or addition of one or several amino acid residues and interacts with a c-Jun protein. (a') a nucleic acid comprising any one of the nucleotide sequences of SEQ ID NOS: 126 to 199, SEQ ID NOS: 200 to 217, SEQ ID NOS: 218 to 224, SEQ ID NOS: 225 to 229, SEQ ID NOS: 230 to 234, SEQ ID NOS: 235 to 238, SEQ ID NOS: 239 to 241, SEQ ID NOS: 242 and 243, SEQ ID NOS: 244 and 245, SEQ ID NOS: 246 and 247, SEQ ID NOS: 248 and 249, SEQ ID NOS: 250 and 251, SEQ ID NOS: 252 and 253, and SEQ ID NOS: 254 and 255, (b') a nucleic acid that hybridizes with a nucleic acid comprising any one of the nucleotide sequences of SEQ ID NOS: 126 to 199, SEQ ID NOS: 200 to 217, SEQ ID NOS: 218 to 224, SEQ ID NOS: 225 to 229, SEQ ID NOS: 230 to 234, SEQ ID NOS: 235 to 238, SEQ ID NOS: 239 to 241, SEQ ID NOS: 242 and 243, SEQ ID NOS: 244 and 245, SEQ ID NOS: 246 and 247, SEQ ID NOS: 248 and 249, SEQ ID NOS: 250 and 251, SEQ ID NOS: 252 and 253, and SEQ ID NOS: 254 and 255, under a stringent condition and encodes a protein that interacts with a c-Jun protein.

- 118. (Previously Presented) The method according to claim 117, wherein the protein comprises any one of the amino acid sequences of SEQ ID NOS: 1 to 69, SEQ ID NOS: 70 to 87, SEQ ID NOS: 88 to 94, SEQ ID NOS: 95 to 99, SEQ ID NOS: 100 to 104, SEQ ID NOS: 105 to 108, SEQ ID NOS: 109 to 111, SEQ ID NOS: 112 and 113, SEQ ID NOS: 114 and 115, SEQ ID NOS: 116 and 117, SEQ ID NOS: 118 and 119, SEQ ID NOS: 120 and 121, SEQ ID NOS: 122 and 123 and SEQ ID NOS: 124 and 125.
- 119. (Previously Presented) The method according to claim 117, wherein the nucleic acid comprises any one of the nucleotide sequences of SEQ ID NOS: 126 to 199, SEQ ID NOS: 200 to 217, SEQ ID NOS: 218 to 224, SEQ ID NOS: 225 to 229, SEQ ID NOS: 230 to 234, SEQ ID NOS: 235 to 238, SEQ ID NOS: 239 to 241, SEQ ID NOS: 242 and 243, SEQ ID NOS: 244 and 245, SEQ ID NOS: 246 and 247, SEQ ID NOS: 248 and 249, SEQ ID NOS: 250 and 251, SEQ ID NOS: 252 and 253, and SEQ ID NOS: 254 and 255.

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120. (Previously Presented) A method for screening for a prey that interacts with a bait, which comprises the step of detecting an interaction between the bait and a prey by the method according to claim 117 and the step of selecting a prey for which an interaction is detected.

- 121. (Previously Presented) A method for screening for a prey that interacts with a bait, which comprises the step of detecting an interaction between the bait and the prey by the method according to claim 118 and the step of selecting a prey for which an interaction is detected.
- 122. (Previously Presented) A method for screening for a prey that interacts with a bait, which comprises the step of detecting an interaction between the bait and a prey by the method according to claim 119 and the step of selecting a prey for which an interaction is detected.